



# Modeling (some aspects of ) the female reproductive system

Romain Yvinec

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Submitted on 19 Jan 2021

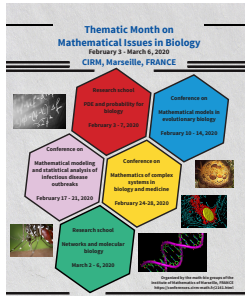
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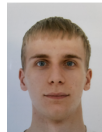
Romain Yvinec

Physiologie de la Reproduction et des Comportements  
INRAe Tours



# Acknowledgements

- ★ INRIA Saclay : Frédérique Clément, Guillaume Ballif, Frédérique Robin
- ★ INRAE PRC : Team BIOS, BINGO (Danielle Monniaux, Véronique Cadoret, Rozenn Dalbies-Tran)
- ★ INRAE LPGP (Julien Bobe, Violette Thermes)
- ★ CEMRACS 2018 (Céline Bonnet (CMAP, X), Kerloun Chahour (U. Côte d'Azur))



# The Mammalian female reproductive system : a complex multiscale system

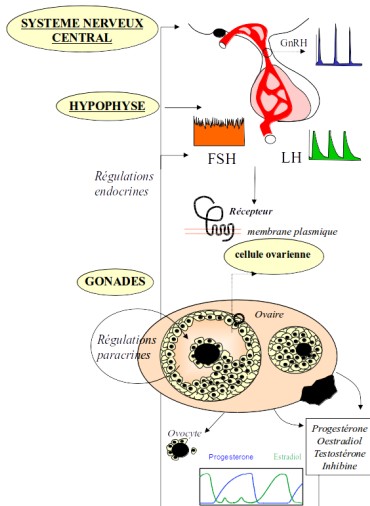
Encoding and decoding **neuro-hormonal signals**

Population dynamics : **gametogenesis**

Intra-cellular level : **signaling networks**



Yvinec et al., *Advances in computational modeling approaches of pituitary gonadotropin signaling*, Expert Opinion on Drug Discovery, 2018.





# Scientific and societal issues in reproductive science

## **Understanding of a complex process of developmental biology, occurring during the whole lifespan**

- Numerous cell types involved, and various interactions
- Many different spatial and temporal scales
- Hormonal feedback (endocrine, paracrine, autocrine)
- Steric and biophysical constraint

## **Preserve the reproductive ability**

- Iatrogenic or physiological alterations
- Sensibility to environmental conditions
- Biodiversity preservation

## **Control of the reproduction function (in humans and animals)**

- Biotechnology of reproduction (*in vivo*, *ex vivo*, *in vitro*)
- Clinical, economical and environmental issues

# The Mammalian female reproductive system : a complex multiscale system

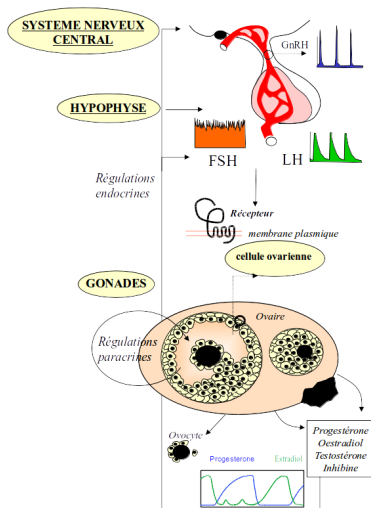
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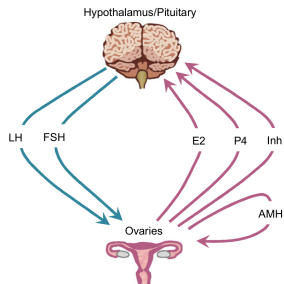


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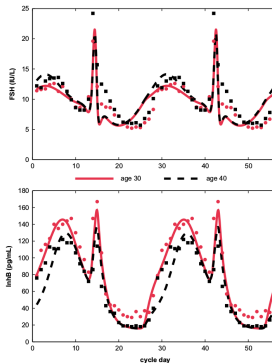


# Neuro-hormonal signals (at the anatomic scale)

Mostly phenomenological equations (DDEs/SDEs) to represent measured levels of circulating hormones, on a daily basis.



*Margolskee & Selgrade, JTB 2013*

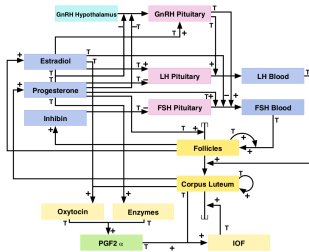


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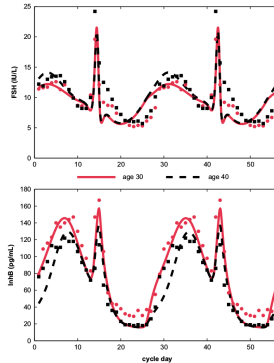
These models can explain some disorders in hormonal levels and predict the effect of pharmaceutical intervention.

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These models can explain some disorders in hormonal levels and predict the effect of pharmaceutical intervention. **Theoretical analysis gets rapidly challenging !**

# The Mammalian female reproductive system : a complex multiscale system

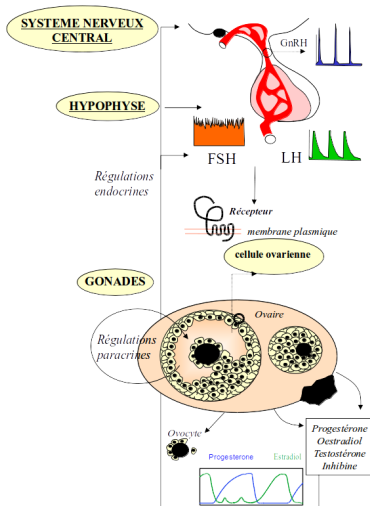
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**Population dynamics : gametogenesis**

Intra-cellular level : **signaling networks**



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# Gametogenesis : Ovarian folliculogenesis

- Morphogenesis and maturation of ovarian follicles  
somatic and germ (egg) cells  
⇒ Somatic cell division and germ cell growth up to ovulation

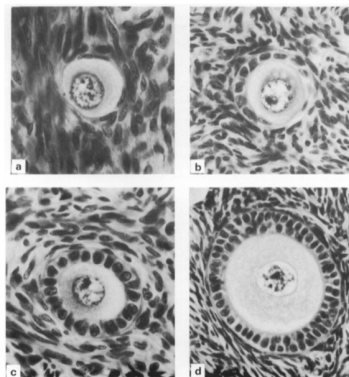
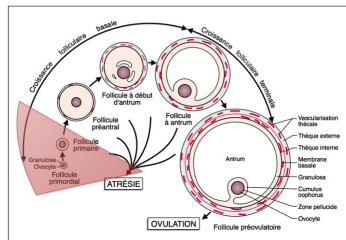


Fig. 1. Illustrations of follicle types: (a) Type B,  $\times 570$ ; (b) Type B/C,  $\times 570$ ; (c) Type C,  $\times 570$ ; (d) Type D,  $\times 410$ .

*Gougeon & Chainy, J. Reprod. Fert. 1987*

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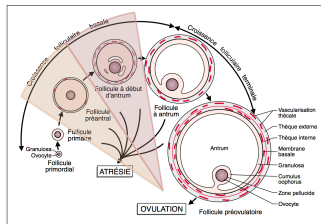
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somatic and germ (egg) cells
- Pool of Quiescent follicles  
static reserve (perinatal in most mammals)  
Slow activation



Monniaux, *Theriogenology* 2016

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Spanning over several ovarian cycles

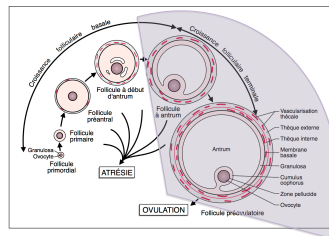


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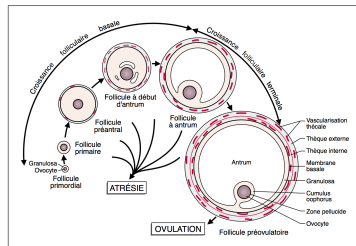
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 Spanning over several ovarian cycles
- Terminal growth  
 After puberty : **ovulation** within an ovarian cycle



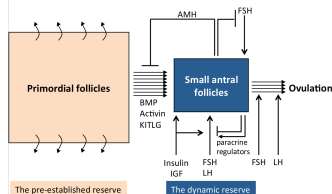
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After puberty : ovulation within an ovarian cycle
- Interactions between all follicles via complex (neuro-) hormonal signals



Ovarian reserves of follicles and their regulations

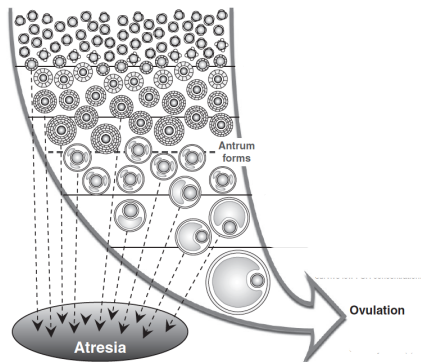


Monniaux, Theriogenology 2016

# Order of magnitude

## Follicle population in women

- Quiescent follicles
  - peri-natal  $\approx 5 \cdot 10^6$
  - At birth  $\approx 1 \cdot 10^6$
  - At puberty  $10^4 - 10^6$
  - At menopause  $< 10^3$
  - Activation rate "A few per days"

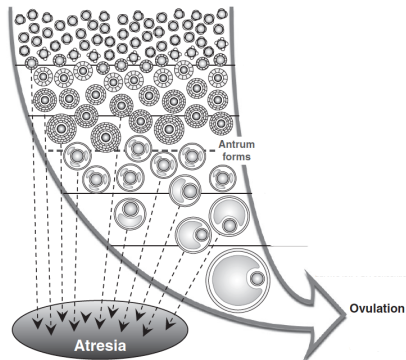


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  - Activation rate "A few per days"
- Growing follicles
  - Maturation time 120 – 180j
  - Basal follicles  $10^3 - 10^4$
  - Terminal follicles  $10^2$
  - Pre-Ovulatory follicles a few
  - Atresia Most of them !



Scaramuzzi et al., *Reprod.Fert. Dev.* 2011

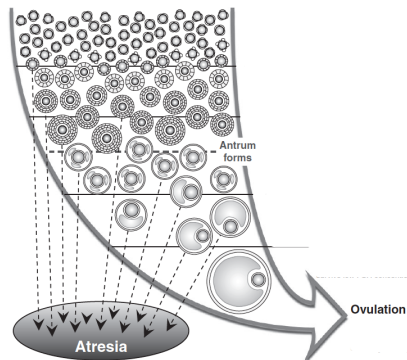
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  - Growing follicles
 

Maturation time	120 – 180j
Basal follicles	$10^3 - 10^4$
Terminal follicles	$10^2$
Pre-Ovulatory follicles	a few
Atresia	Most of them !
- > **Only 400 follicles will ever reach the pre-ovulatory stage**

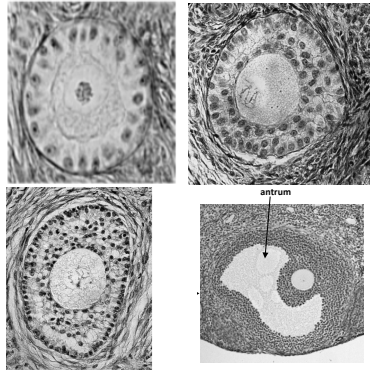


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# Order of magnitude

- **a single follicle (in women)**  
at different maturation stages
 

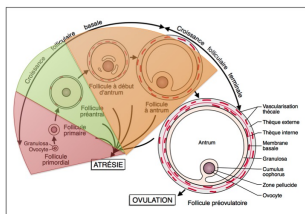
somatic cells diam.	$10\mu m$
ovocyte (egg cell) diam. :	$10 - 100\mu m$
follicle diam.	$0.03 - 20mm$
nb somatic cells	$10^2 - 10^7$



*Courtesy of Danielle Monniaux.*

# Modeling ovarian folliculogenesis

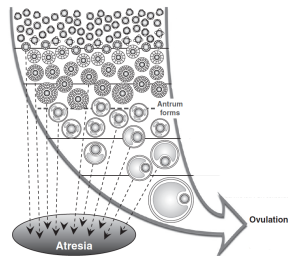
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## Populations of follicles

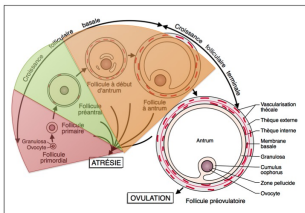


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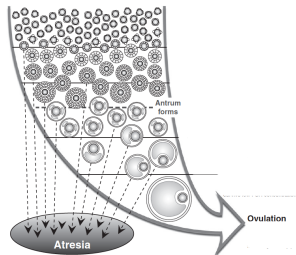
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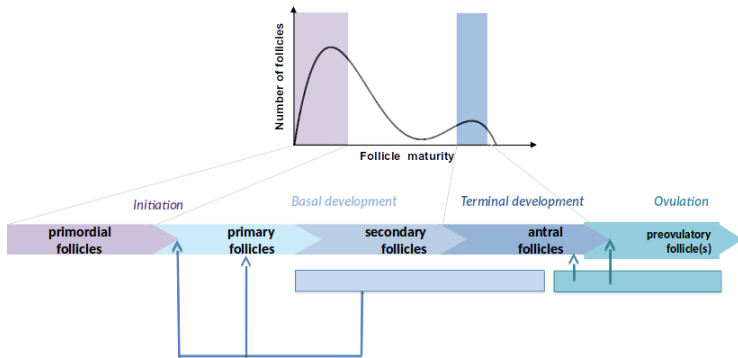
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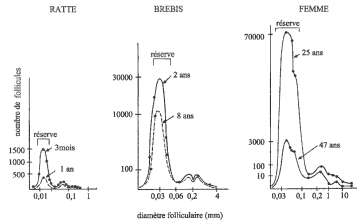
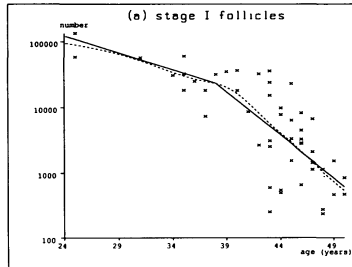
# Models of population of follicles on the lifespan

- ⇒ Nonlinear interactions between follicles populations (endocrine and paracrine)
- ⇒ Slow decay of total follicles number and "stable" repartition in the maturity space

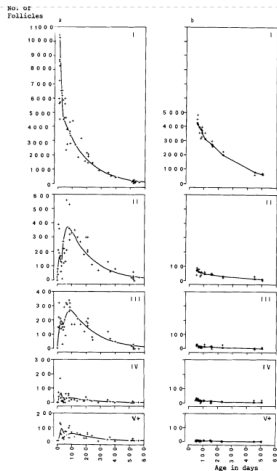


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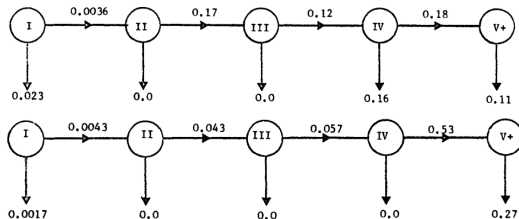
# Models of population of follicles on the lifespan



## An Analytical Model for Ovarian Follicle Dynamics

M. J. FADDY,<sup>1</sup> ESTHER C. JONES<sup>2</sup> AND R. G. EDWARDS<sup>3</sup>  
<sup>1</sup> Department of Mathematical Statistics, University of Birmingham, Birmingham B15 2TT, U.K.; <sup>2</sup> Department of Anatomy, University of Birmingham, Birmingham B15 2TJ, U.K.; and <sup>3</sup> Physiological Laboratory, University of Cambridge, Cambridge CB2 3EG, U.K.

- "Migration-death" stochastic model
- Linear model with time-dependent coefficient



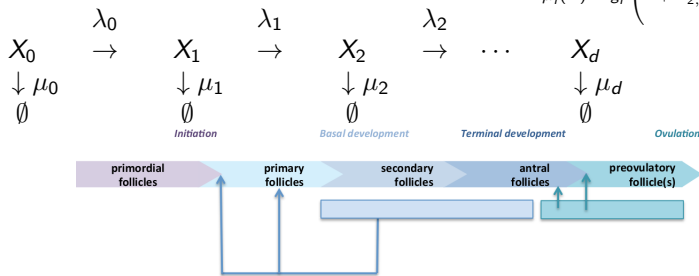
Faddy et al., J. Exp. Zool. 1976

# Models of population of follicles on the lifespan

- Compartment based model (CTMC)
- Nonlinear interaction between follicles populations via  $\lambda$ 's and  $\mu$ 's.
- Several time scales

$$\lambda_i(X) = m_i + \frac{f_i}{1 + K_{1,i} \sum_{j=0}^d \omega_{1,j} X_j},$$

$$\mu_i(X) = g_i \left( 1 + K_{2,i} \sum_{j=0}^d \omega_{2,j} X_j \right)$$



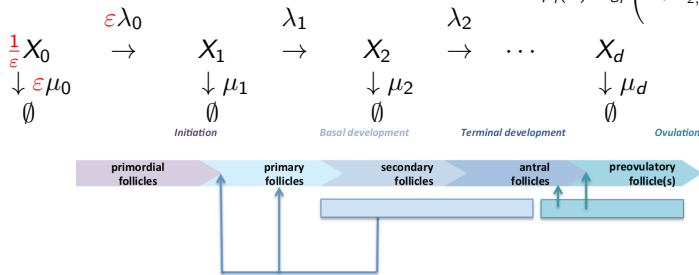
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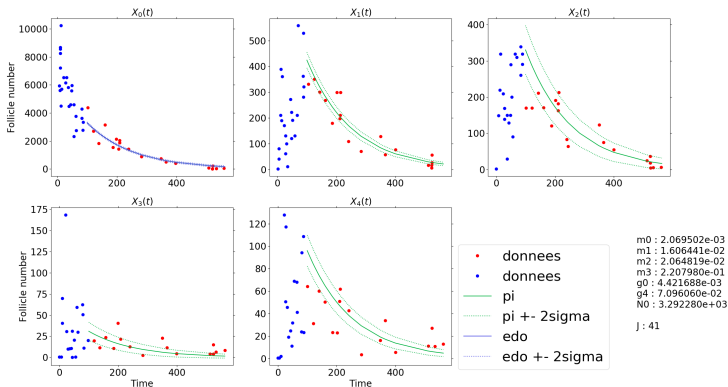
$$\begin{array}{ccccccc}
 \frac{1}{\varepsilon} X_0 & \xrightarrow{\varepsilon \lambda_0} & X_1 & \xrightarrow{\lambda_1} & X_2 & \xrightarrow{\lambda_2} & \dots & X_d \\
 \downarrow \varepsilon \mu_0 & & \downarrow \mu_1 & & \downarrow \mu_2 & & & \downarrow \mu_d \\
 \emptyset & & \emptyset & & \emptyset & & & \emptyset
 \end{array}$$

## Theorem (G. Ballif, Averaging slow-fast dynamics)

For smooth and lower/upper bounded rates  $\lambda, \mu$ , the CTMC converges as  $\varepsilon \rightarrow 0$  (in  $\mathcal{D}_{\mathbb{R}}[0, \infty[ \times \mathcal{L}_m(\mathbb{N}^d)$ ) towards the solution of

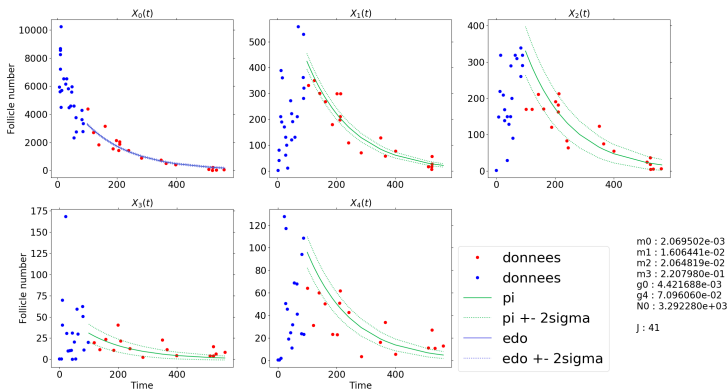
$$\left\{ \begin{array}{ll}
 \frac{dx_0}{dt}(t) &= -\Lambda_0(x_0(t))x_0(t) & x_0(0) = x_0^{in}, \\
 \Lambda_0(x_0) &= \sum_{y \in \mathbb{N}^d} \left( \lambda_0(x_0, y) + \mu_0(x_0, y) \right) \pi_{x_0}(y) & \forall x_0 \in \mathbb{R}_+, \\
 0 &= \sum_{y \in \mathbb{N}^d} L_{x_0} f(y) \pi_{x_0}(y) & \forall f \in \mathcal{C}_b^2(\mathbb{N}^d), \\
 L_{x_0} f(y) &= \lambda_0(x_0, y) x_0 \left[ f(y + e_1) - f(y) \right] & \forall y \in \mathbb{N}^d, \\
 &+ \sum_{i=1}^{d-1} \lambda_i(x_0, y) y_i \left[ f(y + e_{i+1} - e_i) - f(y) \right] \\
 &+ \sum_{i=1}^d \mu_i(x_0, y) y_i \left[ f(y - e_i) - f(y) \right]
 \end{array} \right.$$

# Models of population of follicles on the lifespan



- Separation of time scale is coherent with biological knowledge and follicles count data.

# Models of population of follicles on the lifespan

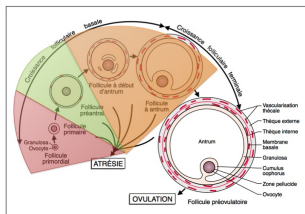


- Separation of time scale is coherent with biological knowledge and follicles count data.
- **Work in progress...**
- Identifiability issues (even with the linear model)
- The limit model allows to test null hypothesis (no regulation)



# Modeling ovarian folliculogenesis

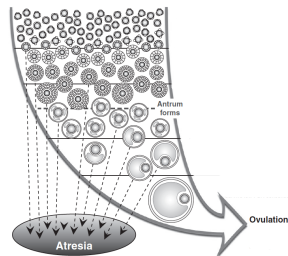
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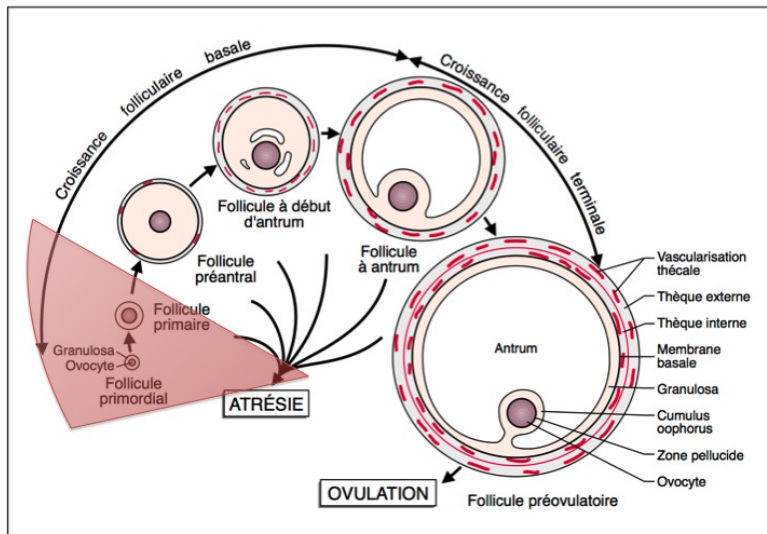
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# Follicle initiation



# Key features of follicle initiation

- Leave the **quiescent** phase (static reserve)
- A single layer of somatic cells
- **Two types** of cells :  
Flattened and Cuboid
- **Irreversible transition** from  
Flattened to Cuboid cells
- The follicle is "activated"  
when all cells have  
transitioned
- **"Awakening" signals both  
from external and internal  
cues**

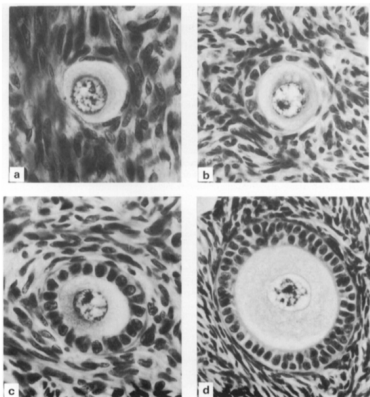
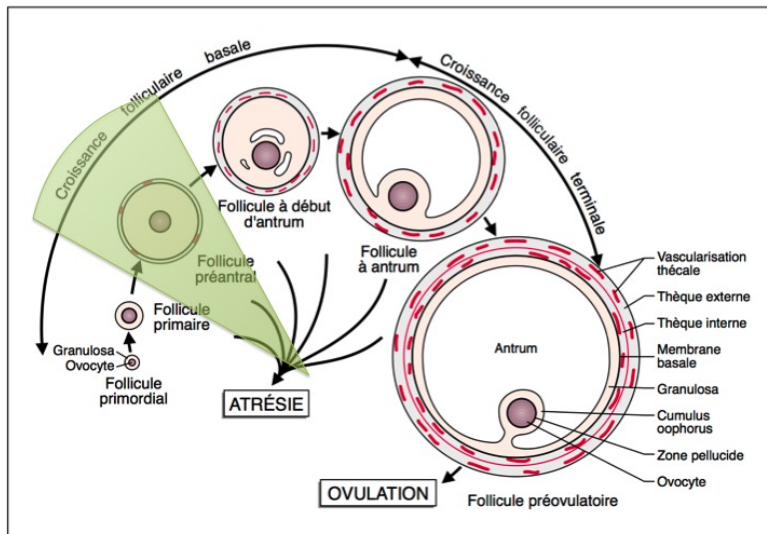


Fig. 1. Illustrations of follicle types: (a) Type B,  $\times 570$ ; (b) Type B/C,  $\times 570$ ; (c) Type C,  $\times 570$ ; (d) Type D,  $\times 410$ .

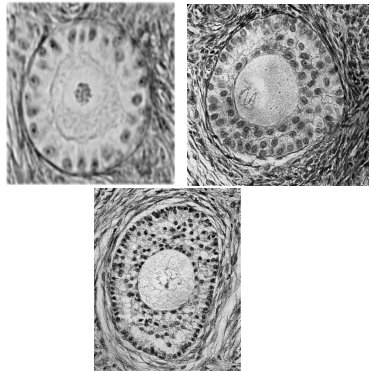
Gougeon & Chainy, *J. Reprod. Fert.* 1987

# Basal growth



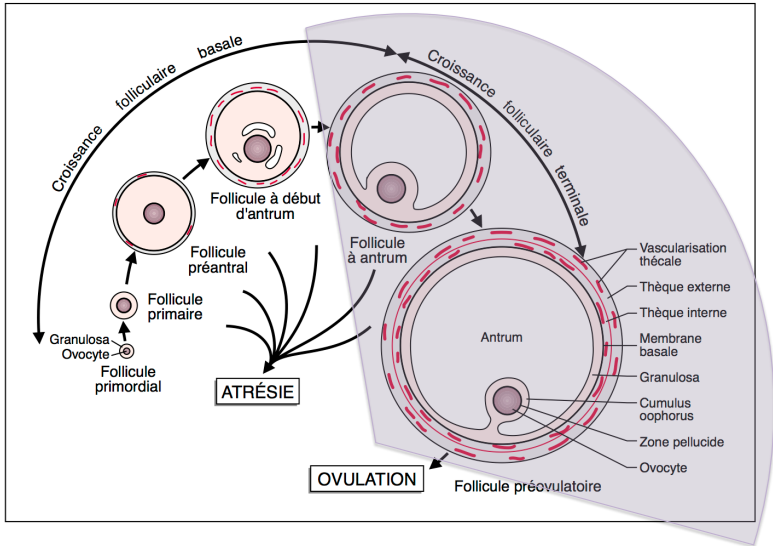
# Key features of follicle basal growth

- Growth of a small follicle after initiation
- **Spherical Symmetry**
- Spatial structure of somatic cells in concentric **layers**
- Joint dynamic
  - ★ Ovocyte **growth**
  - ★ Somatic cells **Proliferation**
- Growth signals from the Ovocyte to somatic cells and *vice-versa*.



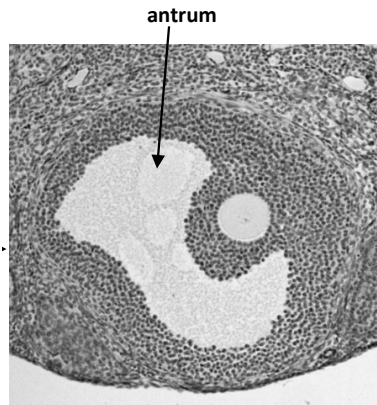
*Courtesy of Danielle Monniaux.*

# Terminal growth



# Key features of Follicle terminal growth

- Lost of spherical symmetry
- Joint Dynamic
  - ★ Liquid-filled cavity formation and growth
  - ★ Switch from proliferation to differentiation of somatic cells
  - ★ Morphogen gradient
- **Role of the Liquid-filled cavity ?**

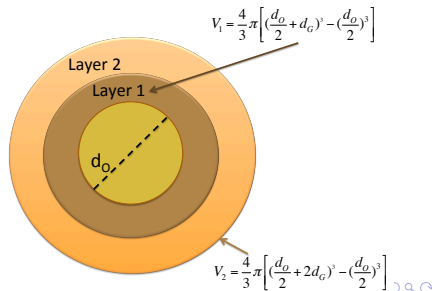
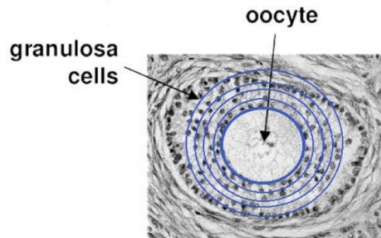


*Tertiary ( antral) follicle*

# Simple model : spatial compartment in successive layers

- Spherical ovocyte ( $d_o$ )
- Linear proliferation dynamics of somatics cells
- Layer dependent division rate
- Multi-type Bellman-Harris Branching process

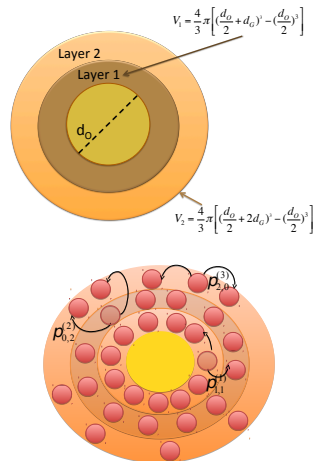
Somatic cells divide and migrate to successive layers.





- The geometrical model allows a simple spatial description
- The model is linear and decomposable : **exponential growth**, with a stable asymptotic spatial profile : there exists a unique  $\lambda > 0$  such that the process  $Z_t$  verifies

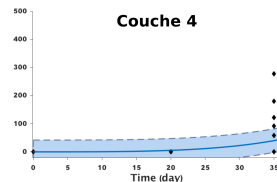
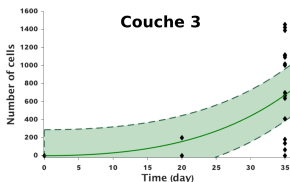
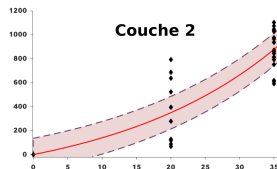
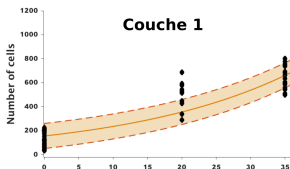
$$\lim_{t \rightarrow \infty} Z_t e^{-\lambda t} = \hat{Z} \quad (\text{in law})$$



Clément et al. *Analysis and Calibration of a Linear Model for Structured Cell Populations with Unidirectional Motion : Application to the Morphogenesis of Ovarian Follicles*, SIAM App. math, 2019

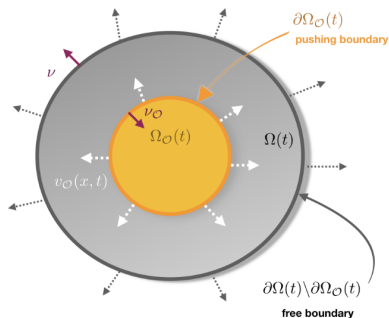
# Fitting results

- ⇒ **Exponential** growth dominated by the first cell layer
- ⇒ Parameter identifiability and doubling time quantification ( $\approx 16$  days) :  
Cell-cycle time ↗ with ovocyte distance



Clément et al. *Analysis and Calibration of a Linear Model for Structured Cell Populations with Unidirectional Motion : Application to the Morphogenesis of Ovarian Follicles*, SIAM App. math, 2019

# More realistic model ?



$\frac{\partial u}{\partial t} + \text{div}(\vec{v}u) = b(x)u(t, x)$   
for  $x \in \Omega(t)$  and with  $\vec{v}$  linked  
to the negative gradient of the  
pressure, and the pressure related  
to the density...

Under locally constant density  
and spherical geometry, one  
have :

$$\frac{d}{dt}(r_F(t)) = \gamma \int_{\Omega(t)} b(x) dx + \frac{d}{dt}(r_O(t))$$

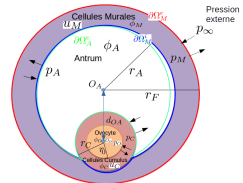
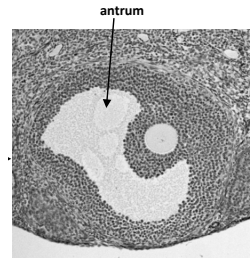
work in progress...

# Antrum growth model

- Multiphasic advection-diffusion-reaction PDE

$$\begin{aligned}
 (\partial_t + \text{div}_x(v_M \cdot)) u_M &= R_M \\
 v_M &= -\mu_M \nabla_x p_M, \\
 p_M &= C_M (u_M - u_0)^{\gamma_M}, \\
 (\partial_t + \text{div}_x(v_a \cdot)) \Phi_a &= D \Delta \Phi_a + R_a \\
 \text{div}_x v_a &= 0, \\
 \nabla \Phi_a \cdot \vec{n}_a &= s(t) = \kappa \int_{\Omega_M} u_M(t, x) dx \\
 \frac{d}{dt} \left( \frac{4}{3} \pi r_a(t)^3 \right) &= J_{H_2O} = L_p(t) (\Delta \Pi - \Delta p) \\
 \Delta \Pi &= c_a \Phi_a^{\gamma_a} \\
 \Delta p &= p_M(t, |x| = r_a^+) - p_e \\
 L_p(t) &= \frac{\Pi(r_F(t) - r_a(t))^4}{8\eta\epsilon(u_M)} n(u_M),
 \end{aligned}$$

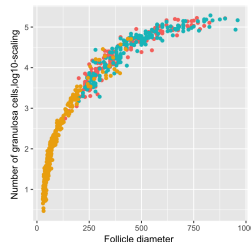
+ Boundary conditions and constitutive laws  
work in progress...



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 \end{aligned}$$



+ Boundary conditions and constitutive laws  
work in progress...

# The Mammalian female reproductive system : a complex multiscale system

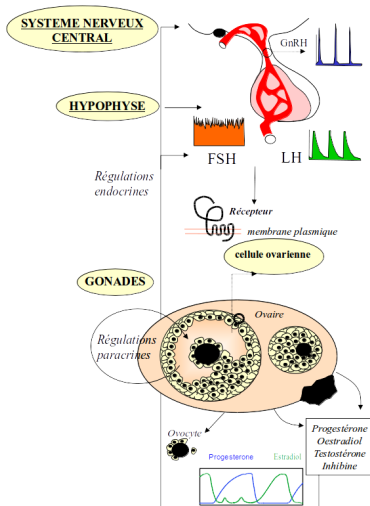
Encoding and decoding **neuro-hormonal signals**

Population dynamics : **gametogenesis**

Intra-cellular level : **signaling networks**



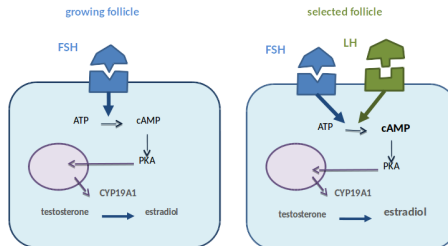
Yvinec et al., *Advances in computational modeling approaches of pituitary gonadotropin signaling*, Expert Opinion on Drug Discovery, 2018.



# cAMP-induced FSH and its regulation at the cellular level

- FSHR signaling network and short-term cAMP induction.
  - Long-term regulation of the FSHR network during follicular selection.
- ⇒ The Cellular "switch" is a pre-requisite for follicle selection .
- ⇒ This switch is implemented at molecular level by the FSHR network and cAMP output (which is a good marker of follicle maturity).

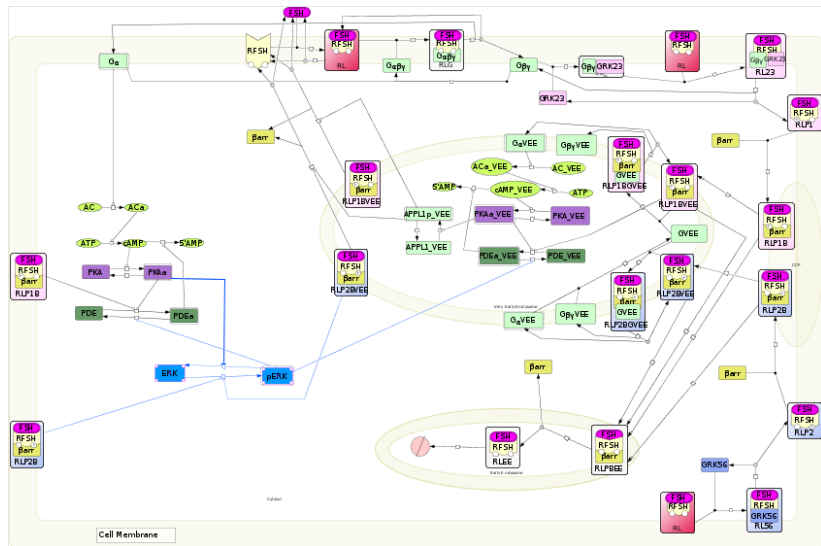
## GRANULOSA CELL







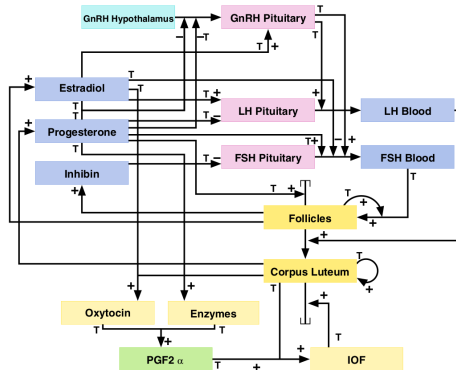
# FSHR networks : very complicated dynamics !



"Network Biology"-> next week !

# Summary

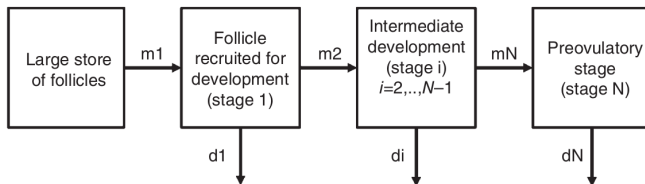
- Circulating hormones dynamical models (Selgrade's model)
  - ★ Purely hormonal dynamics
  - ★ limit cycles in compact delay differential equation models of the ovarian cycle



Stötzel et al., *Therio*. 2012

# Summary

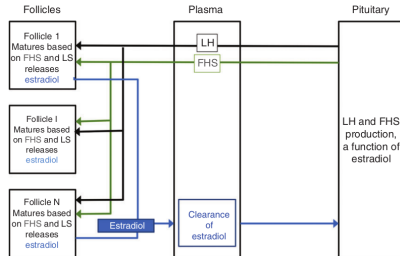
- Circulating hormones dynamical models (Selgrade's model)
- Follicles dynamics in compartment (Faddy's model)
  - ★ Reproductive lifespan timescale
  - ★ Follicle communication and "hormonal control" through population feedback



Clark & Kruger, WIREs Syst Biol Med 2017

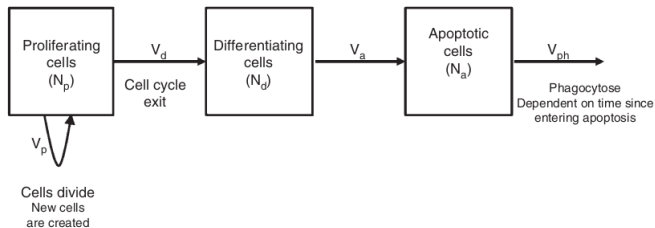
# Summary

- Circulating hormones dynamical models (Selgrade's model)
- Follicles dynamics in compartment (Faddy's model)
- Coupled Hormonal/Follicle dynamical models (Lacker's & Clément's model)
  - ★ Ovarian cycle / follicle cohort timescale
  - ★ Follicle cohort subject to a shared hormonal environment
  - ★ The individual follicle maturity rate has local positive feedback and global negative feedback.



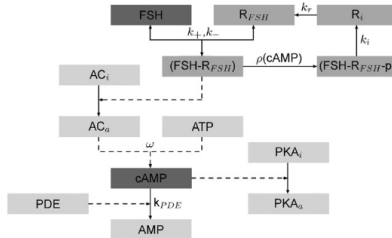
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- Circulating hormones dynamical models (Selgrade's model)
- Follicles dynamics in compartment (Faddy's model)
- Coupled Hormonal/Follicle dynamical models (Lacker's & Clément's model)
- Cell dynamics in a single follicle (Clément's model)
  - ★ Cell cycle time scale
  - ★ Pool of different cell types within a single follicle
  - ★ Complex geometry and moving boundary problems



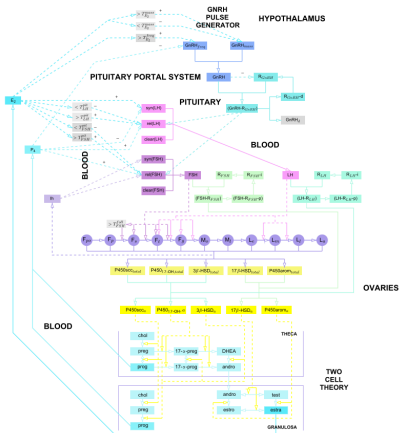
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- Coupled Hormonal/Follicle dynamical models (Lacker's & Clément's model)
- Cell dynamics in a single follicle (Clément's model)
- Intra-cellular dynamics (Clément / Quignot & Bois)
  - ★ Steroidogenesis, cAMP response



# Multiscale Models of ovarian follicle selection

- Putting "many" pieces together (Reinecke & Deuflhard)



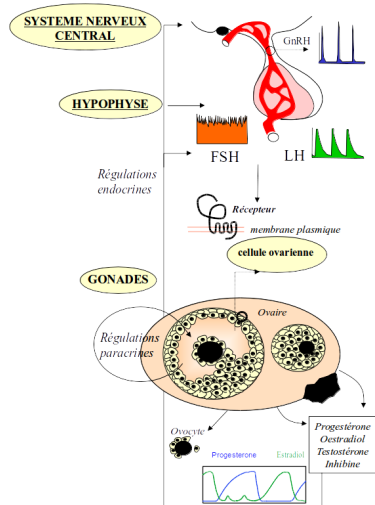
- ★ From (stochastic) GnRH pulse generator to detailed steroid metabolism through (compartment-based) follicle development
- ★ *Focus in this paper is on the model development*
- ★ 43 equations (stochastic input and delay differential equations), 191 parameters.

Reinecke & Deuflhard, JTB 2007

# Summary

## Population dynamics in ovarian folliculogenesis

- Somatic cells proliferation, differentiation, and migration during follicle initiation & growth
- Multiscale nonlinear dynamics shape the follicle population distribution into different maturity stages.

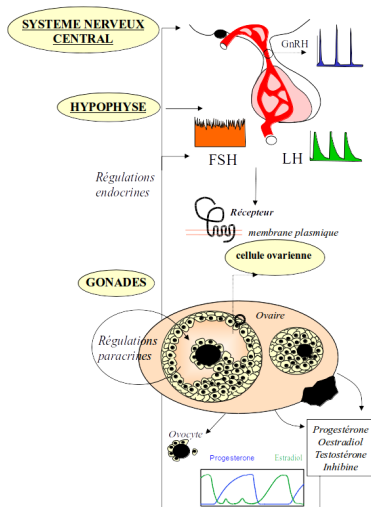




# Summary

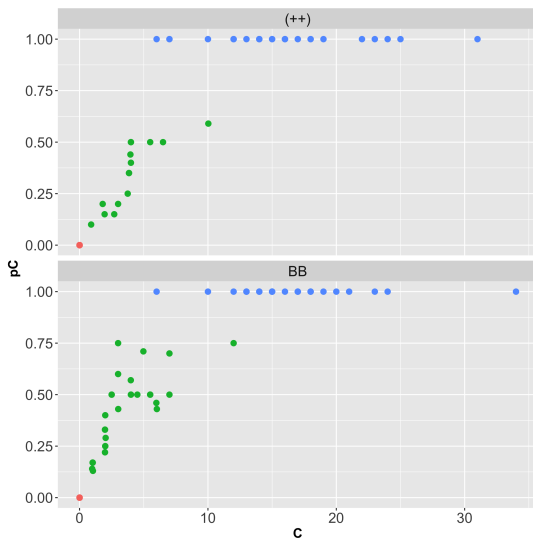
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Thank you for your attention !

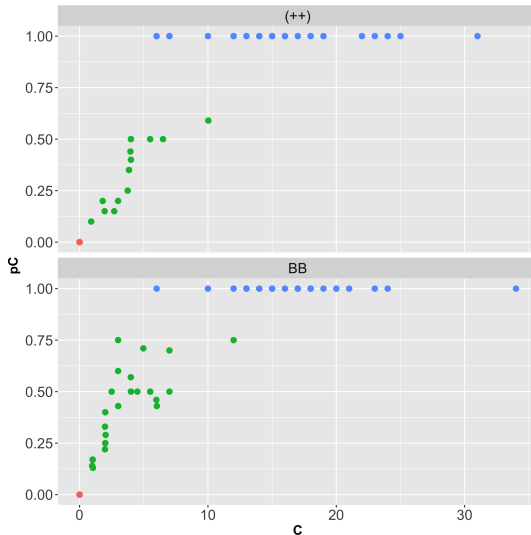
# Ex vivo data (snapshot data)



- Ex vivo data in sheep fetus ( Courtesy of K. McNatty) : WT (++) vs Mutant (BB)

⇒ **Proportion of cuboid cells**  
 $p_C = C / (F + C)$  vs  
**number of cuboid cells  $C$**

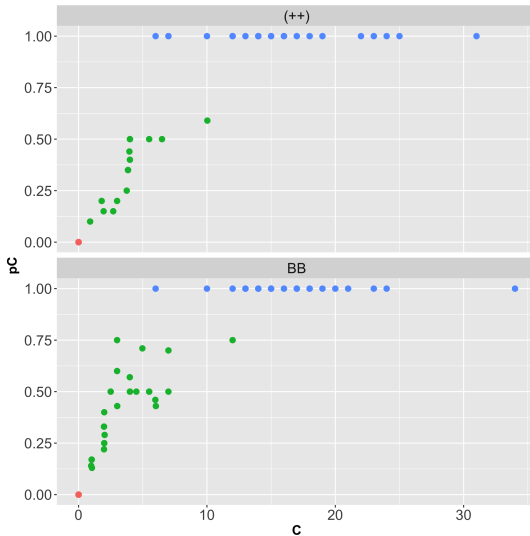
# Ex vivo data (snapshot data)



- Ex vivo data in sheep fetus WT vs BB
- Once activated, follicles have "fast" cell proliferation

⇒ **Are both differentiation et de proliferation process concomitant or successive ?**

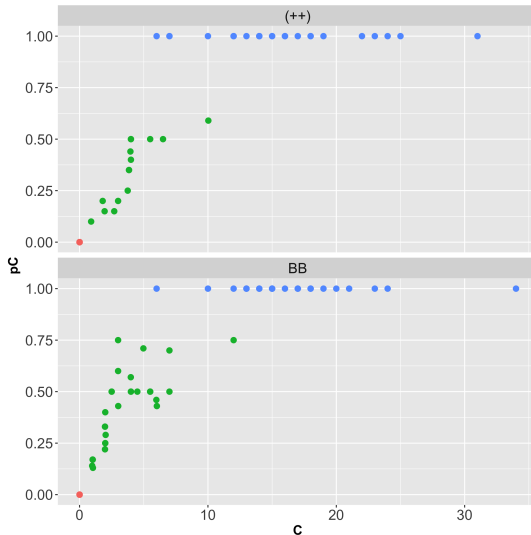
# Ex vivo data (snapshot data)



- Ex vivo data in sheep fetus WT vs BB
- Proportion of cuboid cells seems higher in mutant than WT, for a given number of cuboid cells.

⇒ **Is it coming from a kinetic difference ?**

# Ex vivo data (snapshot data)



- Ex vivo data in sheep fetus WT vs BB
- Regulatory mechanism for this process are barely known.

⇒ **Is the transition of cell differentiation abrupt or more progressive ?**

Events	Reaction	Intensity function
differentiation	$F \rightarrow C$	$\alpha F + \beta \frac{FC}{F + C}$
prolifération	$C \rightarrow C + C$	$\gamma C$

- ↔ Two cell populations :  $F$  (flattened) and  $C$  (cuboid)
- ↔ Small number of cells
- ↔ Retro-action of cuboid cells on the differentiation rate : is it relevant ?
- ↔ From  $(F_0, 0)$  to  $(0, C_T)$

- Theoretical study

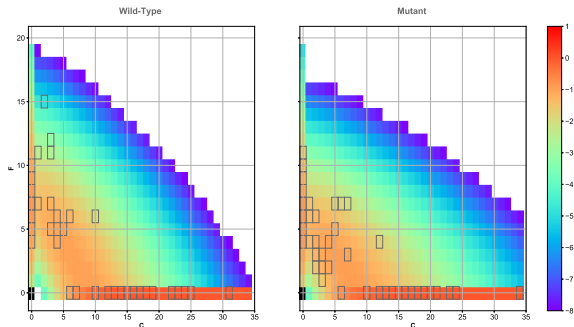
- ⇒ Statistics of the "transition" time  $\tau$  to reach  $F = 0$ .
- ⇒ Variability of final cuboid cells ( $\mathbb{E}[C_\tau] < \infty$  if  $\gamma < \alpha + \beta$ )
- ⇒ Impact of parameters e.g. on qualitative dynamics (progressive vs abrupt)

- Parameter calibration : lack of identifiability. Either  $\gamma \ll 1$  and  $\beta$  unconstrained, or  $\gamma > 1$  and  $\beta/\gamma \gg 1$



Robin et al. *Stochastic nonlinear model for somatic cell population dynamics during ovarian follicle activation*, (submitted) arXiv :1903.01316

# Agreement to data



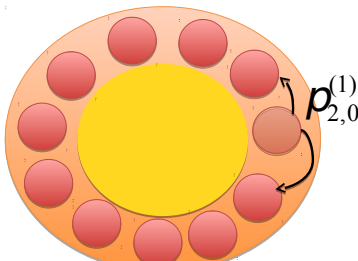
- ⇒ The model can capture both data sets
- ⇒ Lack of identifiability (non-conclusive on retro-action)
- ⇒ First differentiation, then proliferation (slightly more concomitant in mutant case)



Robin et al. *Stochastic nonlinear model for somatic cell population dynamics during ovarian follicle activation*, (submitted) arXiv :1903.01316

# Dynamical model (Multi-type Bellman-Harris Branching process)

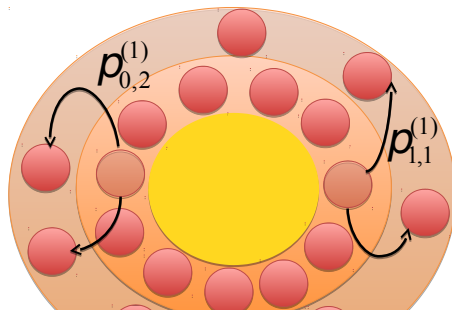
- **Age** and **position** dependent division rate (cell cycle regulated by the ovocyte)
- At division, unidirectional motion **centrifugal**
- Cells are **independant** between each other (Unlimited layer capacity)





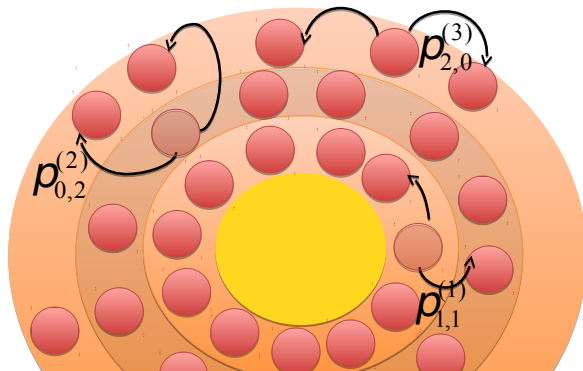
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# Data

- We have counting data of somatic cells in snapshot data, morphological data (diameter) and *order of magnitude of transit times between follicle "type"*

	$t = 0$	$t = 20$	$t = 35$
#Data points	34	10	18
Total cell number	$113.89 \pm 57.76$	$885.75 \pm 380.89$	$2241.75 \pm 786.26$
Oocyte diameter ( $\mu m$ )	$49.31 \pm 8.15$	$75.94 \pm 10.89$	$88.08 \pm 7.43$
Follicle diameter ( $\mu m$ )	$71.68 \pm 13.36$	$141.59 \pm 17.11$	$195.36 \pm 23.95$

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⇒ Can we explain proliferation in concentric layers by a simple model of "division-migration"? Or do physical constraint play important role?

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- ⇒ Can we characterize the growth rate of a follicle and spatial repartition of somatic cells ?
- ⇒ What is the impact of spatial position of a somatic cell on its division rate ?